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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Database :

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SUMMARIES

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RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
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RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Mount S.M., Nelson K.A., Nixon K., Nusskern D.R., Paeleb J.M.,
RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
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RA Sylers B.C., Stapleton M., Skupski M.P., Smith T.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Zheng X.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Yao Q.A.,
RA Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Zhu X., Smith H.O.,
RA Zheng X.H., Zhong F.N., Rubin G.M., Venter J.C.,
RT "The genome sequence of Drosophila melanogaster.",
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CC -i-TISSUE Specificity: TeSTIS.
CC -i-Developmental Stage: PRIMARY SPERMATOCYTES.
CC -i-Dowaln: THIS PROTEIN IS MOSTLY COMPOSED OF REBETITIVE C-G-P
the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement.
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Amanatides P.G., Scherer S.E., Li P.W.,
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ch. Dev. 35:143-151(1991).
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DISULFID
SEQUENCE
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          Eumalacostraca; Eucarida;
                    Scylla serrata (Mud crab).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
                                                                                                                                                                                                                                                                                                      Serine protease inhibitor; DISULFID 3 36 DISULFID 12 32 DISULFID 16 28
                                                                                                                                                                                                                                                                                                                                                                                  "Primary structure and properties of the cathepsin G/chymotrypsin inhibitor from the larval hemolymph of Apis mellifera."; Eur. J. Blochem. 262:680-687(1999).
-i- FUNCTION: CHYMOTRYPSIN AND CATHEPSIN G INHIBITOR.
-i- SUBCELLULAR LOCATION: SECRETED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Apis mellifera (Honeyhee).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
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15-JUL-1999 (Rel. 38, Last sequence up
30-MAY-2000 (Rel. 39, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Spermatogenesis; Repeat; Multigene family. SEQUENCE 55 AA; 5225 MW; 95A12F3AEC88BB6C CRC64;
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Portunoidea;
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Portunidae;
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SEQUENCE.
                                                                                01-APR-1993 (Rel. 25, Created)
01-NOV-1997 (Rel. 35, Last sequence up
01-NOV-1997 (Rel. 35, Last annotation
METALLOTHIONEIN-1 (CUMT-1).
Homarus americanus (American lobster).
Eukaryota; Metazoa; Arthropoda; Crusta
Eumalacostraca; Eucarida; Decapoda; Pl
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                             MEDLINE-89215793; PubMed=2709004; Brouwer M., Winge D.R., Gray W.R.; "Structural and functional diversi
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                                                     TISSUE=HEPATOPANCREAS;
                                                            SEQUENCE
                                                                           Nephropoidea;
                                                                                                                                         P29499;
                                                                                                                                                MT1_HOMAM
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nes 5; Conser
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                                                                                                                                                                                                    ructural and functional diversity of American lobster Homarus americanus.
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               Biochem.
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                                                                           Nephropidae;
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                                                                                                                                                STANDARD;
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                                                                                          Crustacea; Malacostraca;
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                                                                                 Pleocyemata;
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Best Local
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J. Biol. Chem. 257:2420-2426(1982).

-i- FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT RESIDUES THAT BIND VARIOUS HEAVY METALS. CLASS
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Biochemistry 33:8858-8865(1994).
                                                                                                                                                                                                                                            MT1_SCYSE STANDARD; PRT; 58 AA. P02805; P02805
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Lerch K., Ammer D., Olafson R.W.;
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                                                                                                 "Crab metallothionein.
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HSSP; P55949; 1DMF.
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IPR003019;
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gnments and
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INTERPRO; IPRO02045;
INTERPRO; IPRO03019;
PFAM; PF00131; metalth
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P55950;
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Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca; Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyu Eubrachyura; Portunoidea; Portunidae; Callinectes.
                                                                                                                                                                                                                                                                                                                       01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
01-NOV-1997 (Rel. 35, Last annotation update)
METALLOTHIONEIN-II (MT-II) (MT-IIB/MT-IIA).
                                                                                                                                                                                                       Brouwer M., Enghild J., Hoexum-Brouwer T., Thogersen I., 'Primary structure and tissue-specific expression of blue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  METAL
SEQUENCE
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                                                                                INTERPRO; IPRO03019; PFAM; PF00131; metalt
                                                                                                                                                             (Callinectes sapidus) metallothionein isoforms.";
Biochem. J. 311:617-622(1995).
-i- FUNCTION: BINDS SIX DIVALENT METAL IONS. KNOW
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S; PR00858; MTCRUSTACEAN.
-binding; Metal-thiolate cluster; C.
N 1 29 BETA.
N 30 58 ALPHA.
N 30 58 CLUSTER B.
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                                                    Chelation; Copper;
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e crab
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Metal-binding;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Astacus fluviatilis (Broad-fingered crayfish) (Astacus astacus). Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca; Eucarida; Decapoda; Pleocyemata; Astacidea;
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-1- MASS SPECTROMETRY: MW=5910.8;
-1- SIMILARITY: BELONGS TO FAMILY
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-!- INDUCTION: BY CADMIUM.
-!- SIMILARITY: BELONGS TO
HSSP; P55949; 1DMD.
INTERPRO; IPR002045; -.
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"Purification and characterization of a cadmium-induced metallothionein from the shore crab Carcinus maenas (L.)
Biochem. J. 297:609-614(1994).
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Carcinus maenas (Common shore crab) (Green crab).

Curcinus maenas (Common shore crab) (Green crab).

Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;

Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;

Eubrachyura; Portunoidea; Portunidae; Carcinus.
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RESIDUES THAT BIND VARIOUS HEA
CRUSTACEA ARE INVOLVED IN THE
OF HEAVY-METAL IONS.
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22; Mismatches
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INTERPRO;
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1. FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTE RESIDUES THAT BIND VARIOUS HEAVY METALS. CLASS I MTS IN CARE INVOLVED IN THE SEQUESTRATION OF ELEVATED LEVELS OF THE SECONDARY OF THE SECONDARY OF THE SECONDARY METALS.
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Eumalacostraca; Eucarida; Decapo
Eubrachyura; Portunoidea; Geryon
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01-NOV-1997 (Rel. 35,
METALLOTHIONEIN (MT).
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-1- MASS SPECTROMETRY: MW=6156.8; METHOD=MALDI.
-1- SIMILARITY: BELONGS TO FAMILY 3 IN METALLOTHIONEIN SUPERFAMILY
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                                                                                                                                                                                                                                                                                                                      TISSUE=MIDGUT
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Decapoda; Pleocyemata; Brachyu
Geryonidae; Potamon.
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PDB; 1DMF; 07-FEB-95.
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Marula S.S., Brouwer M., Hua Y., Armitage I.M.;
"Three-dimensional solution structure of Callinectes sapidus metallothionein-1 determined by homonuclear and heteronuclear magnetic resonance spectroscopy.";
Biochemistry 34:620-631(1995).
-i- FUNCTION: BINDS SIX DIVALENT METAL IONS. KNOWN TO BIND CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca; Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura; Eubrachyura; Portunoidea; Portunidae; Callinectes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=96033062; PubMed=7487904; Brouwer M., Enghild J., Hoexum-Br
                                                                                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PR0085
Metal-binding;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Callinectes
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Pred. No.
4398B0A9D1A96D34 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           cluster;
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                                                           (IN MT-IA)
                                                                                                                                                                                                                                                                                                                                                                                                                                         Chelation; Copper; Cadmium;
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                                                                                                                                                                                                                                                                                                  Harris N.L., Harvey D., Helman T.J., Hernandez J.R., Houck J.,
A Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
A Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
A Jalali M., Kalush F., Karpen G.H., Ke Z., Kunnison J.A., Ketchum K.A.,
A Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
A Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
A Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.A.,
A Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
A Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
A Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
A Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
A Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
A Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
A Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
A Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
Spradling A.C., Stapleton M., Strong R., Sun E.,
Syriskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 5
                                    Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J.S., Zhen, Zhang G., Zhao Q., Zheng L., Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Zhu X., Smith H.O., Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.; "The genome sequence of Drosophila melanogaster.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beallew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottler P., Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottler P., Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I. Cherry J.M., Cawley S., Dahke C., Davenport L.B., Davies P., Cherry J.M., Cawley S., Dahke C., Davenport L.B., Davies P., de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M., Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn F. Dourbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W. Cabrielista E.C., Ferraz C., Ferriera S., Fleischmann W. Cabrielista E.C., Ferraz C., Ferriera S., Fleischmann W. Cabrielista C.C., Ferraz C., Ferra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K., Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F., George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "A cluster of four genes selectively expressed in the male of Drosophila melanogaster.";
Mech. Dev. 35:143-151(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       M84A_DROME STANDARD; PRT; 001642; O9VIA3; 01-JUL-1993 (Rel. 26, Created) 01-JUL-1993 (Rel. 26, Last sequence up 01-OCT-2000 (Rel. 40, Last annotation MALE SPECIFIC SPERM PROTEIN MST84DA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. STRAIN-BERKELEY;
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nce 287:2185-2195(2000).
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Boersch D., Glaetzer K.H., Schaefer
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RESULT 13
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01-JUL-1993
01-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota;
Pterygota;
FLYBASE; FBgn0004175; Mst84Dd.
Spermatogenesis; Repeat; Multigene family.
SEQUENCE 68 AA; 6480 MW; 2F2BD26128DE3DEF CRC64;
                                                                                                                                                          modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                        of Drosophila melanogaster.";
Mech. Dev. 35:143-151(1991).
-i- TISSUE SPECIFICITY: TESTIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        "A cluster of four genes selectively of Drosophila melanogaster.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Spermatogenesis; Repeat; Multigene family. SEQUENCE 63 AA; 5806 MW; BF84CD74CBCEFD3F CRC64;
                                                                                                EMBL; X67703; CAA47940.1;
                                                                                                                                          or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                               the European Bioinformatics Institute.
                                                                                                                                                                                                                                                      between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schaefer M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=92102953; PubMed=1684716;
Kuhn R.., Kuhn C., Boersch D., Gla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=OREGON-R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ephydroidea;
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DOMAIN: THIS PROTEIN IS MOSTLY COMPOSED OF REPETITIVE C-G-P
                                                                                                                                                                                                                                                                                                                     SIMILARITY: BELONGS TO THE MST(3)CGP FAMILY
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DEVELOPMENTAL STAGE: PRIMARY SPERMATOCYTES
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                                                                                                                                                                                                        non-profit institutions as long
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Neoptera; Endopterygo
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                                                                                                                                                            license agreement (See http://www.isb-sib.ch/announce/
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ilidae; Drosophila
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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams M.D., Celniker S.E., Holt R.A., Asbburner M., Henderson S.N.,
RA George R.A., Lewis S.E., Richards S., Asbburner M., Henderson S.N.,
RA George R.A., Lewis S.E., Richards S., Asbburner M., Henderson S.N.,
RA George R.A., Lewis S.E., Richards S., Asbburner M., Henderson S.N.,
RA Burting G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Burting M.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfelifer B.D.,
RA Ballew R.M., Bayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,
RA Ballew R.M., Bayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dun P.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dun P.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Ketchum K.A.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegyam C.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Llang Y., Lin X.,
RA Herkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Herkulov G., Milshina N.Y., Mobarry C., Morris J., Moshrefi A.,
RA Herkulov G., Milshina N.Y., Mobarry C., Morris J., Moshrefi A.,
RA Herkulov G., Siden K.A., Nixon K., Nussekrn D.R., Pacleb J.M.,
RA Herkulov G., Siden K.A., Nixon K., Nussekrn D.R., Smith T.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Syler E., Spradling A.C., Stapleton M., Strong S., Yao Q.A.,
RA Hard G., Chang S., Pan S., Pollard J., Wang S.,
RA Wang K.H., W
               RESULT 14
A62F_DROME
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Best Local S
Matches 5
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046202;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=20196006; PubMed=10731132;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=CANTON-S; TISSUE=MALE ACCESSORY GLAND MEDLINE=98135120; PubMed=9474779;
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Pterygota;
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01-OCT-2000
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5; Conserv
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Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Metazoa; Padontervqota; Diptera; Brachycera; Muscomorpha;
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18.5%;
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Pred. No. 26;
22; Mismatches
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Science 287:2185-2195(2000).
-!- FUNCTION: RESPONSIBLE FOR PHYSIOLOGICAL AND BEHAVIORAL CHANGES IN

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RESULT 1.
VE5_RHPV1
ID VE5_RHPV1
P24834;
P24834;
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Matches 5; Conserv
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01-MAR-1992
01-OCT-1996
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SEQUENCE
                                                                                                                               Early protein.
SEQUENCE 157 AA;
                                                                                                                                                          EMBL; M60184; AAA79315.1; ALT_SEQ. EMBL; M60184; AAA79316.1; ALT_SEQ. PIR; F38503; W5WLR1.
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                                                                                                                                                                                                                                                              the European Bioinformatics Institute. The use by non-profit institutions as long
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                                                                                                                                                                                                                                                                                                                                    Virology 181:424-429(1991).
                                                                                                                                                                                                                                                                                                                                                Ostrow R.S., Labresh K.V., Faras A.J.; "Characterization of the complete RhPV 1 genomic sequence and integration locus from a metastatic tumor.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; U85763; AAB96387.1; -. EMBL; AE003475; AAF47683.1; -. FLYBASE; FBgn0020509; Acp62F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Viruses;
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Viruses; dsDNA viruses, no RNA stage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41 TECPVACPETCEYSGNGPCVKMCGAPC 67
 23
                          SUBCELLULAR LOCATION: SECRETED (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MATED FEMALE FLIES
VLCKANCCCYACKPPPFCCFWLCFCCC 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        115 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Rel. 21, Created)
(Rel. 21, Last sequence update)
(Rel. 34, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
                                                                                                                                17398 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12570 MW;
                                                                       88.2%;
18.5%;
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18.5%; Pred. No.
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Pred. No.
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                                                                                                                                AC7AA67158844686 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     157 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                     Papovaviridae; Papillomavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            update)
                                                                       DB 1;
46;
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37;
                                                                                                                                                                                                                                                           There are no restrictions on ong as its content is in no
                                                                                                                                                                                                                                               Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                        0;
                                                                                   Length 157;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 115
                                                          Indels
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Search completed: March 1, Job time: 399 sec

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